DT05 Rec'd PCT/PTO 0 3 JAN 2005

SEQUENCE LISTING

<110> MARTIN, Annette

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SANGAR, DAVID V.
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     RIJNBRAND, RENE
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			gtg Val													3284
_			acg Thr 985			_	_		_		_		_			3332
	Gly		ccc Pro			Ala					Glu					3380
Pro	_	_	gga Gly	_	Val		_			Arg	_	_				3428
	Ala		gcc Ala	Gln					Leu					Ile		3476

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ggc tgg ccc gc Gly Trp Pro Al 1110			_	_	-
ggc tcc tcg ga Gly Ser Ser As		Leu Val Thr			Pro
gtg cgc cgg cg Val Arg Arg Ar 114	g Gly Asp				
att tcc tac tt Ile Ser Tyr Le 1160	-		Gly Pro Leu		
gga cac gcc gt Gly His Ala Va 1175	al Gly Leu				
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agc ttc cag gt Ser Phe Gln Va 122	al Ala His				
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ctc aac ccc to Leu Asn Pro Se 1255	er Val Ala				
aag gcc cat gg Lys Ala His Gl 1270			Arg Thr Gly	Val Arg Thr	Ile
	1275		1280		1285

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1530 1535 1540

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2505 2510 2515

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gcc tgc tac tcc ata Ala Cys Tyr Ser Ile 2870				96
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. . . .

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9599

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<213> Hepatitis C virus

<400> 14

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Leu Leu Arg Ile Pro Gln Ala Ile Met Asp Met Ile Ala Gly Ala His Trp Gly Val Leu Ala Gly Ile Ala Tyr Phe Ser Met Val Gly Asn Trp Ala Lys Val Leu Val Val Leu Leu Phe Ala Gly Val Asp Ala Glu Thr His Val Thr Gly Gly Asn Ala Gly Arg Thr Thr Ala Gly Leu Val Gly Leu Leu Thr Pro Gly Ala Lys Gln Asn Ile Gln Leu Ile Asn Thr Asn Gly Ser Trp His Ile Asn Ser Thr Ala Leu Asn Cys Asn Glu Ser Leu Asn Thr Gly Trp Leu Ala Gly Leu Phe Tyr Gln His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Arg Leu Thr Asp Phe Ala Gln Gly Trp Gly Pro Ile Ser Tyr Ala Asn Gly Ser Gly Leu Asp Glu Arg Pro Tyr Cys Trp His Tyr Pro Pro Arg Pro Cys Gly Ile Val Pro Ala Lys Ser Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val Val Gly Thr Thr Asp Arg Ser Gly Ala Pro Thr Tyr Ser Trp Gly Ala Asn Asp Thr Asp Val Phe Val Leu Asn Asn Thr Arg Pro Pro Leu Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Ser Thr Gly Phe Thr Lys Val Cys Gly Ala Pro Pro Cys Val Ile Gly Gly Val Gly Asn Asn Thr Leu Leu Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr Tyr Ser Arg Cys Gly Ser Gly Pro Trp Ile Thr Pro Arg Cys Met Val Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Ile Asn Tyr Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Leu Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Ser Thr Thr Gln Trp Gln Val Leu Pro Cys Ser Phe Thr Thr Leu Pro Ala Leu Ser Thr Gly Leu Ile His Leu His Gln Asn Ile Val Asp Val Gln Tyr Leu Tyr Gly Val Gly Ser Ser Ile Ala Ser Trp Ala Ile Lys Trp Glu Tyr Val Val Leu Leu Phe Leu Leu Ala Asp Ala Arg Val Cys Ser Cys Leu Trp Met Met Leu Leu Ile Ser Gln Ala Glu Ala Ala Leu Glu Asn Leu Val Ile Leu Asn Ala Ala Ser Leu Ala Gly Thr His Gly Leu Val Ser Phe Leu Val Phe Phe Cys Phe Ala Trp Tyr Leu Lys Gly Arg Trp Val Pro Gly Ala Val Tyr Ala Leu Tyr Gly Met Trp Pro Leu Leu Leu Leu Leu Leu Ala Leu Pro Gln Arg Ala Tyr Ala Leu Asp Thr Glu Val Ala Ala Ser Cys Gly Gly Val Val Leu Val Gly Leu Met Ala Leu Thr Leu Ser

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Phe	Leu 850	Thr	Arg	Val	Glu	Ala 855	Gln	Leu	His	Val	Trp 860	Val	Pro	Pro	Leu
Asn 865	Val	Arg	Gly	Gly	Arg 870	Asp	Ala	Val	Ile	Leu 875	Leu	Met	Cys	Val	Val 880
	Pro	Thr	Leu	Val 885		Asp	Ile	Thr	Lys 890		Leu	Leu	Ala	Ile 895	
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Val	Arg	Val 915		Gly	Leu	Leu	Arg 920		Cys	Ala	Leu	Ala 925		Lys	Ile
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Gly	Cys	Ile		Thr 1045	Ser	Leu	Thr		Arg L050	Asp	Lys	Asn		Val 1055	Glu
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Сув	Ile	Asn 1075	Gly	Val	Cys	-	Thr 1080	Val	Tyr	His		Ala L085	Gly	Thr	Arg
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Thr	Pro	Cys		Cys 1125	Gly	Ser	Ser	_	Leu 1130	Tyr	Leu	Val		Arg 1135	His
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_	Val			1285				:	1290				:	1295	
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Met Ser Ala Asp Leu Glu Val Val Thr Ser Thr Trp Val Leu Val Gly
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Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Thr Gly Cys Val
1665 1670 1675
Val Ile Val Gly Arg Ile Val Leu Ser Gly Lys Pro Ala Ile Ile Pro
          1685 1690 1695
Asp Arg Glu Val Leu Tyr Gln Glu Phe Asp Glu Met Glu Glu Cys Ser
                      1705
Gln His Leu Pro Tyr Ile Glu Gln Gly Met Met Leu Ala Glu Gln Phe
                   1720
                                 1725
Lys Gln Lys Ala Leu Gly Leu Leu Gln Thr Ala Ser Arg His Ala Glu
                1735
                               1740
Val Ile Thr Pro Ala Val Gln Thr Asn Trp Gln Lys Leu Glu Val Phe
1745 1750 1755
Trp Ala Lys His Met Trp Asn Phe Ile Ser Gly Ile Gln Tyr Leu Ala
          1765 1770 1775
Gly Leu Ser Thr Leu Pro Gly Asn Pro Ala Ile Ala Ser Leu Met Ala
                      1785
Phe Thr Ala Ala Val Thr Ser Pro Leu Thr Thr Gly Gln Thr Leu Leu
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1795	1800	1805	
Phe Asn Ile Leu Gly 1810	Gly Trp Val Ala 1815	Ala Gln Leu Ala Ala 1 1820	Pro Gly
	Val Gly Ala Gly 1830	Leu Ala Gly Ala Ala 3	Ile Gly 1840
	Lys Val Leu Val	Asp Ile Leu Ala Gly	
		Phe Lys Ile Met Ser (
Val Pro Ser Thr Glu 1875		Leu Leu Pro Ala Ile I 1885	Leu Ser
Pro Gly Ala Leu Val 1890	Val Gly Val Val 1895	Cys Ala Ala Ile Leu A	Arg Arg
	Glu Gly Ala Val 1910	Gln Trp Met Asn Arg 1 1915	Leu Ile 1920
Ala Phe Ala Ser Arg 1925		Ser Pro Thr His Tyr V 1930 19	Val Pro 935
1940	· 1945	Ala Ile Leu Ser Ser I 1950	
1955	1960	Gln Trp Ile Ser Ser (
1970	1975	Arg Asp Ile Trp Asp 7	_
1985	1990	Trp Leu Lys Ala Lys I	2000
2005			015
2020	2025	His Thr Arg Cys His (2030	
2035	2040	Gly Thr Met Arg Ile V 2045	-
2050	2055	Gly Thr Phe Pro Ile 2	
2065	2070	Pro Ala Pro Asn Tyr I 2075	2080
2085			095
2100	2105	Thr Thr Asp Asn Leu 1 2110	
2115	2120	Phe Thr Glu Leu Asp (2125	
2130	2135	Lys Pro Leu Leu Arg (2140	
3	Gly Leu His Glu 2150	Tyr Pro Val Gly Ser (31n Leu 2160
	Pro Asp Val Ala	Val Leu Thr Ser Met	
		Ala Gly Arg Arg Leu 2	
Gly Ser Pro Pro Ser 2195	Met Ala Ser Ser 2200	Ser Ala Ser Gln Leu S 2205	Ser Ala
2210	2215	Asn His Asp Ser Pro 2220	
2225	2230	Arg Gln Glu Met Gly (2235	2240
2245			255
2260	2265	Arg Glu Val Ser Val 1 2270	
Glu Ile Leu Arg Lys 2275	Ser Arg Arg Phe 2280	Ala Arg Ala Leu Pro V 2285	Jal Trp

Ala Arg Pro Asp Tyr Asn Pro Pro Leu Val Glu Thr Trp Lys Lys Pro 2295 Asp Tyr Glu Pro Pro Val Val His Gly Cys Pro Leu Pro Pro Pro Arg 2310 2315 Ser Pro Pro Val Pro Pro Pro Arg Lys Lys Arg Thr Val Val Leu Thr 2325 2330 2335 Glu Ser Thr Leu Ser Thr Ala Leu Ala Glu Leu Ala Thr Lys Ser Phe 2340 2345 2350 Gly Ser Ser Ser Thr Ser Gly Ile Thr Gly Asp Asn Thr Thr Thr Ser 2355 2360 2365 Ser Glu Pro Ala Pro Ser Gly Cys Pro Pro Asp Ser Asp Val Glu Ser 2380 2370 2375 Tyr Ser Ser Met Pro Pro Leu Glu Gly Glu Pro Gly Asp Pro Asp Leu 2385 2390 2395 Ser Asp Gly Ser Trp Ser Thr Val Ser Ser Gly Ala Asp Thr Glu Asp 2405 2410 2415 Val Val Cys Cys Ser Met Ser Tyr Ser Trp Thr Gly Ala Leu Val Thr 2420 2425 Pro Cys Ala Ala Glu Glu Gln Lys Leu Pro Ile Asn Ala Leu Ser Asn 2435 2440 2445 Ser Leu Leu Arg His His Asn Leu Val Tyr Ser Thr Thr Ser Arg Ser 2455 2460 Ala Cys Gln Arg Gln Lys Lys Val Thr Phe Asp Arg Leu Gln Val Leu 2465 2470 2475 2480 Asp Ser His Tyr Gln Asp Val Leu Lys Glu Val Lys Ala Ala Ala Ser 2485 2490 2495 Lys Val Lys Ala Asn Leu Leu Ser Val Glu Glu Ala Cys Ser Leu Thr 2500 2505 2510 Pro Pro His Ser Ala Lys Ser Lys Phe Gly Tyr Gly Ala Lys Asp Val 2515 2520 2525 Arg Cys His Ala Arg Lys Ala Val Ala His Ile Asn Ser Val Trp Lys 2530 2535 2540 Asp Leu Leu Glu Asp Ser Val Thr Pro Ile Asp Thr Thr Ile Met Ala 2545 2550 2555 Lys Asn Glu Val Phe Cys Val Gln Pro Glu Lys Gly Gly Arg Lys Pro 2565 2570 2575 Ala Arg Leu Ile Val Phe Pro Asp Leu Gly Val Arg Val Cys Glu Lys 2580 2585 2590 Met Ala Leu Tyr Asp Val Val Ser Lys Leu Pro Leu Ala Val Met Gly 2595 2600 2605 Ser Ser Tyr Gly Phe Gln Tyr Ser Pro Gly Gln Arg Val Glu Phe Leu 2615 2620 Val Gln Ala Trp Lys Ser Lys Lys Thr Pro Met Gly Phe Ser Tyr Asp 2625 2630 2635 Thr Arg Cys Phe Asp Ser Thr Val Thr Glu Ser Asp Ile Arg Thr Glu 2645 2650 2655 Glu Ala Ile Tyr Gln Cys Cys Asp Leu Asp Pro Gln Ala Arg Val Ala 2665 2670 Ile Lys Ser Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Leu Thr Asn 2680 2685 Ser Arg Gly Glu Asn Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val 2695 2700 Leu Thr Thr Ser Cys Gly Asn Thr Leu Thr Cys Tyr Ile Lys Ala Arg 2705 2710 2715 Ala Ala Cys Arg Ala Ala Gly Leu Gln Asp Cys Thr Met Leu Val Cys 2725 2730 Gly Asp Asp Leu Val Val Ile Cys Glu Ser Ala Gly Val Gln Glu Asp 2745 Ala Ala Ser Leu Arg Ala Phe Thr Glu Ala Met Thr Arg Tyr Ser Ala 2760 2765 Pro Pro Gly Asp Pro Pro Gln Pro Glu Tyr Asp Leu Glu Leu Ile Thr

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	Cys	Ser	Ser		Val 2790	Ser	Val	Ala		Asp 2795	Gly	Ala	Gly		Arg 2800	
Val	Tyr	Tyr		Thr 2805	Arg	Asp	Pro		Thr 2810	Pro	Leu	Ala	_	Ala 2815	Ala	
Trp	Glu	Thr	Ala 2820	Arg	His	Thr		Val 2825	Asn	Ser	Trp		Gly 2830	Asn	Ile	
Ile		Phe 2835	Ala	Pro	Thr		Trp 2840	Ala	Arg	Met		Leu 2845	Met	Thr	His	
	Phe 850	Ser	Val	Leu		Ala 2855	Arg	Asp	Gln		Glu 2860	Gln	Ala	Leu	Asn	
Cys 2865		Ile	Tyr	_	Ala 2870	Cys	Tyr	Ser		Glu 2875	Pro	Leu	Asp		Pro 2880	
Pro	Ile	Ile		Arg 2885	Leu	His	Gly		Ser 2890	Ala	Phe	Ser		His 2895	Ser	
Tyr	Ser	Pro 2	Gly 2900	Glu	Ile	Asn	-	Val 2905	Ala	Ala	Cys		Arg 2910	Lys	Leu	
Gly		Pro 2915	Pro	Leu	Arg		Trp 2920	Arg	His	Arg		Arg 2925	Ser	Val	Arg	
2	930	Leu			2	2935				:	2940					
2945	;	Asn	_	2	2950	_		_	2	2955				2	2960	
Ala	Ala	Gly		Leu 2965	Asp	Leu	Ser		Trp 2970	Phe	Thr	Ala		Tyr 2975	Ser	
	_		2980	_			2	2985			_	2	2990			
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	> D	escr: rime:	-	on of	E Art	cific	cial	Seq	uence	e: :	Syntl	hetio	С			
<400		6 caa c	7022	raac!	-											19